

## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.**

Application Serial Number: 10/526,108

Source: PCT/10

Date Processed by STIC: 3/16/05

# ***ENTERED***



PCT

## RAW SEQUENCE LISTING

DATE: 03/16/2005

PATENT APPLICATION: US/10/526,108

TIME: 08:27:20

Input Set : N:\Rasheed\PTO.SR.txt

Output Set: N:\CRF4\03162005\J526108.raw

3 <110> APPLICANT: Maier, Sabine  
 5 <120> TITLE OF INVENTION: METHOD AND NUCLEIC ACIDS FOR THE ANALYSIS OF BREAST CELL  
 PROLIFERATIVE  
 6 DISORDERS  
 8 <130> FILE REFERENCE: 47675-102  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/526,108  
 C--> 11 <141> CURRENT FILING DATE: 2005-02-28  
 13 <150> PRIOR APPLICATION NUMBER: PCT/EP2003/007827  
 14 <151> PRIOR FILING DATE: 2003-07-18  
 16 <150> PRIOR APPLICATION NUMBER: DE10255104.9  
 17 <151> PRIOR FILING DATE: 2002-11-26  
 19 <150> PRIOR APPLICATION NUMBER: DE10239313.3  
 20 <151> PRIOR FILING DATE: 2002-08-27  
 22 <160> NUMBER OF SEQ ID NOS: 396

## ERRORED SEQUENCES

**Does Not Comply**  
**Corrected Diskette Needed**

25313 <210> SEQ ID NO: 396  
 25314 <211> LENGTH: 21  
 25315 <212> TYPE: DNA  
 25316 <213> ORGANISM: Artificial Sequence  
 25318 <220> FEATURE:  
 25319 <223> OTHER INFORMATION: CCND2 primer  
 25321 <400> SEQUENCE: 396  
 25323 cctaacctcc ttcctttaac t  
 E--> 25329 - 2 -

21

**VERIFICATION SUMMARY**

DATE: 03/16/2005

PATENT APPLICATION: US/10/526,108

TIME: 08:27:23

Input Set : N:\Rasheed\PTO.SR.txt

Output Set: N:\CRF4\03162005\J526108.raw

L:10 M:270 C: Current Application Number differs, Replaced Application Number  
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:1314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:3300  
L:3152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49 after pos.:1080  
M:341 Repeated in SeqNo=49  
L:10007 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:150 after pos.:3300  
L:10033 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:151 after pos.:240  
L:10790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:162 after pos.:3300  
L:10816 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:163 after pos.:240  
L:17567 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:262 after pos.:1080  
M:341 Repeated in SeqNo=262  
L:17619 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:263 after pos.:720  
M:341 Repeated in SeqNo=263  
L:19179 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:280 after pos.:1080  
M:341 Repeated in SeqNo=280  
L:19231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:281 after pos.:720  
M:341 Repeated in SeqNo=281  
L:25329 M:254 E: No. of Bases conflict, this line has no nucleotides.